OM nucleic - nucleic search, using sw model

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

Run on:

Scoring table: Sequence: Perfect score:

IDENTITY_NUC Gapext 1.0

383533 seqs, 122816752 residues

US-09-676-436-3 4990

Searched:

Post-processing: Minimum Match 0%

Issued_Patents_NA:* Maximum Match 100% Listing first 45 s

summaries

/cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/packfiles1.seq:*

Result

Score

Length DB

SUMMARIES

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0 0

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24 24 24 23.8 23.6

US-08-400-256-4
US-08-975-365-4
US-08-877-097-5
US-08-878-957-5
US-08-878-957-5
US-08-878-957-7
US-08-973-124-78
US-08-973-124-78
PCT-US9-100-664A-5
US-08-100-256-9
US-08-975-365-9
US-08-182-175A-54
US-08-612-158A-8
US-08-672-158A-8
US-08-672-158A-8
US-08-684-704A-17
US-08-684-704A-17
US-08-684-704A-17
US-08-684-704A-17
US-08-684-704A-17
US-08-324-888A-8
US-08-484-324-17
US-08-332-766A-41

0 0

23.8 22.8 22.8 22.6 22.4 22.4 22.4 22.4 22.4

Minimum DB seq length: 0 Maximum DB seq length: 100

Total number of hits satisfying chosen parameters:

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Sequence 5, Appli
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Sequence 79, Appli
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Sequence 63, Appli
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                                                                   sequence 5, Appli
sequence 10, Appl
sequence 21, Appl
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Sequence
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17, Appl
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; patent No. 575049
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US-08-400-256-4/c
                                                                                                                                                          ; MOLECULE TYPE: DNA
US-08-400-256-4
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                                           Query Match
Best Local S
Matches 50
                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 33,728
REGISTRATION NUMBER: 3985
REGISTRATION INFORMATION:
TELEPHONE: 212-867-0123
TELEPHONE: 212-87-0655
TELEPAX: 212-878-9655
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Havelund, Svend
APPLICANT: Halstrom, John
APPLICANT: John
APPLICANT: Andersen, Asser Sloth
APPLICANT: Markussen, Jan
APPLICANT: Markussen, Mar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
No. 57504970 No. 5750497disk of No. 5750497th America, Inc.
ADDRESSEE: No. 57504970 No. 5750497disk of No. 5750497th America, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TIPE: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/400,256
FILING DATE: 03-MAR-1995
FILING DATE: 03-MAR-1995
CLASSIFICATION: 514
ACTIONERY/AGENT INFORMATION:
NAME: Lambiris, Elias J-78
REGISTRATION NUMBER: 33-78
REGISTRATION NUMBER: 33-78
REGISTRATION NUMBER: 34-786,220-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
4469 aagaatgaagootagtagaatatggaottggaaaattotottaatcactactgtatgtaa 4528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 405 Lex
CITY: New York
STATE: New York
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                                                                                                                                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
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                                                      50;
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                                                                         Similarity
                                                                                                                                                                                                                                                           nucleic acid
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                                                                                                                                                                                                                   linear
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                                                                              55.6%;
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US-08-465-594A-75
US-08-93-124-260
PCT-US96-08014-260
PCT-US96-09014-260
PCT-US96-09472-99
US-08-481-697-166
US-08-483-366-514
US-09-243-367-5
US-08-453-463B-10
US-08-445-4648-10
US-08-445-4648-10
US-08-445-4648-10
US-08-413-118-113
US-08-413-118-113
                                                        score 26; DB 1; Pred. No. 6.1e+02; 0; Mismatches 40
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                                                                                                                 Length 100;
                                                                         Indels
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Sequence 260, App
Sequence 260, App
Sequence 99, Appl
Sequence 99, Appl
Sequence 99, Appl
Sequence 31, Appl
Sequence 514, Appl
Sequence 514, Appl
Sequence 55, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 23, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 23, Appl
Sequence 113, Appl
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US-08-427-097-5
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US-08-975-365-4
                                  Sequence 5, Application US/08427097 Patent No. 5668294
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-975-365-4/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 100 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/400
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Elias J
REGISTRATION NUMBER: 33,728
REFERENCE/DOCKET NUMBER: 3985
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
TELEPAX: 212-87-9655
INFORMATION DOS 675
                                                                                                                                                                                                   4529 tatttacataaagactgtgctgagaagcag 4558
                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/08975365 Patent No. 6011007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM CC COmpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                           37 CAATTGGAAAACTACTGTGCTTAGACGCAG 8
                                                                                                                                                                                                                                                          97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTTGTAC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: ACYLATED INSULIN
NUMBER OF SEQUENCES: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37 CAATTGGAAAACTACTGTGCTTAGACGCAG 8
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                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jonassen, Ib
Andersen, Asser Sloth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Havelund, Svend
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                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                       Score 26; DB 3; Length 100;
Pred. No. 6.Te+02;
0; Mismatches 40; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3985.220-US
                                                                                                                                                                                                                                                                                                                                                            40; Indels
                                                                                                                                                                                                                                                                                                                                                            0; Gaps
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                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/08878957
Patent No. 5965796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                 TITLE OF INVENTION: Metal Resistance : TITLE OF INVENTION: Transgenic Plants NUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 80303
ZIP: 80303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE

                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3086 ttgattttctacaagcaattgaacctgcctttat 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (303) 499-8089
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3026 atgtootgaotaaatgtgagagtggtagaggtacaagacccaggtgggcgaotcaaggat 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATER
OPERATING SYSTEM: PC-DO
SOFTWARE: PATENTIN Rele.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 21-APR-199
                                               CITY: Boulder
STATE: Colora
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 99 base pairs
TYPE: nucleic acid
                       COUNTRY:
                                                          ADDRESSEE: Greeniee, wither and STREET: 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL:
ZIP: 80303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 40
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 GTGAGCATACTCAAGCAAGCCAAGTTGCCTATAT 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTTCGCATGGAAGGCATTGAAGTGC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 51; Conserv
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TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRRSSEE: Greenlee, Winner and Sullivan, P.C. STREET: 5370 Manhattan Circle, Suite 201
                                       Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boulder
                                                                                                                                                                                              Meagher, Richard B.
Summers, Anne O.
Rugh, Clayton L.
IVENTION: Metal Resistance Sequences and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           င္ပ
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                                                                                                 Greenlee, Winner and Sullivan,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (303) 499-8080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.5%; Score 25.2; DB 1; Length 99; 54.3%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anne O.
Metal Resistance Sequences and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Transgenic Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US/08/427,097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    33,878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  40-94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
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Qγ В

RESULT

В

Matches

MEDIUM TYPE:

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US-08-458-423A-78/c
US-08-458-423A-78/c
; Sequence 78, Application US/08458423A
; Patent No. 5731144
; PENERAL INFORMATION:
; CENERAL INFORMATION:
; PENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
ETILING DATE: 19-JUN-1997
CLASSIFICATION: 800
PRIOR APPLICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 40-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 499-8089
TELEFAX: (303) 499-8089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/427,097
EILING DATE: 21-APR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 99 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear nucleic acid MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "Oligonucleotide" HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3026 atgtcctgactaaatgtgagagtggtagaggtacaagacccaggtgggcgactcaaggat 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3086 ttgattttctacaagcaattgaacctgcctttat 3119
                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: HIGH AFFINITY TGFA NUCLEIC TITLE OF INVENTION: ACID LIGANDS AND INHIBITORS NUMBER OF SEQUENCES: 89
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDERSTECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/458,423A
FILING DATE: 2-UNNE-1995
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62 GTGAGCATACTCAAGCCAAGCCAAGTTGCCTATAT 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTTCGCATGGAAGGCATTGAAGTGC 61
                                                                                                                                                                         COUNTRY: U
ZIP: 80111
                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 0.5%;
Similarity 54.3%;
51; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
                                                                                                                                                                                                           Denver
: Colorado
                                                                                                                                                                                                                                              E: Swanson and Bratschun, L.L.C.
8400 East Prentice Avenue, Suite #200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NO
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                                                                                                                                                                                                                                                                                                                                                                                  STEVEN RINGQUIST
                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                   LARRY GOLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                      Diskette, 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US/08/878,957
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 25.2;
Pred. No. 1e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pB 2; Length 99;
1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-458-423A-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/117,991
APPLICATION DATE: 8-SEPTEMBER-1993
PRIOR APPLICATION DATE: 1992
APPLICATION NUMBER: 07/931,473
APPLICATION NUMBER: 07/931,473
APPLICATION NUMBER: 07/931,473
APPLICATION NUMBER: 07/931,473
APPLICATION NUMBER: 08/931,473
APPLICATION NUMBER: 08/960
AREGISTRATION NUMBER: 08/960
REGISTRATION NUMBER: 08/960
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 10/93-333
TELETAX: (303) 793-3433
INFORMATION FOR SEQ ID NO: 78:
INFORMATION FOR SEQ ID NO: 78:
INFORMATION FOR SEQ ID NO: 78:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-458-424B-78/c
US-08-458-424B-78/c
Sequence 78, Application US/08458424B
patent No. 5731424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,131
FILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA: 07/536,428
APPLICATION NUMBER: 07/536,428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA: 77964,624
APPLICATION NUMBER: 07/964,624
FILING DATE: 21-OCTOBER-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                        TITLE OF INVENTION: HIGH AFFINITY TGFA NUCLEIC TITLE OF INVENTION: ACID LIGANDS AND INHIBITORS NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: PENELOPE J. TOOTHMAN
APPLICANT: STEVEN RINGQUIST
APPLICANT: LARRY GOLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: single
                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,131
FILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y Match 0.5%; Score 24.6; DB 1; Length 71; Local Similarity 65.5%; Pred. No. 1.2e+03; hes 36; Conservative 0; Mismatches 19; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                            CITY: Denver
                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                            colorado
                                                                                                                                                                                                                                                                                                                                                                                E: Swanson and Bratschun, L.L.C.
8400 East Prentice Avenue, Suite #200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                             USA
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; ANTI-SENSE: US-08-878-957-5

Query Match Best Local : Matches

FILING DATE:

11-JUNE-1990

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US-08-973-124-78/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6207816
GENERAL INFORMATION:
APPLICANT: LARRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 78, Application US/08973124 Patent No. 6207816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                          FILING DATE: 02-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/458,424
                  PRIOR APPLICATION DATA:
                                                                                                                                     CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                          FILING DATE: 30-MAY-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/458,423
FILING DATE: 02-JUNE-1995
                                                                                                                                                                                                                  SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LARRY GOLD et al.
TITLE OF INVENTION: HIGH AFFINITY OLIGONUCLEOTIDE
TITLE OF INVENTION: LIGANDS TO GROWTH
TITLE OF INVENTION: FACTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3958 gaacatgtgattaggctgtattcaaagcagatcaccattgcgatcaacgtcctcc 4012
                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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    APPLICATION NUMBER:
                                    FILING DATE:
                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                               MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Swanson & Blackern, John STREET: 8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 33,960
REFERENCE/DOCKET NUMBER: NED
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,4
FILING DATE: 17-AUGUST-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/964,67
FILING DATE: 21-OCTOBER-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        56 GAAGACGTAAGTAAGTATTAAAAACATACCCCCCCTGGCCGCATCGTCCTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: Si
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Diane H. McCle
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 8-SEPT
                                                                                                                                                                                                                                                                                                                                                                                                                  OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                         Colorado
                                                                                                                                                                                                                                                                                                                            USA
                                                                                                                                                                                                                                                             IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (303)
                             02-JUNE-1995
                                                                                                                                                                                                                                                                            Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       H. McClearn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8-SEPTEMBER-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              793-3433
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                                                                                                                                                                                                                                                MS-DOS
08/465,594
                                                                                                                                     PCT/US96/08014
                                                                                                                                                                                               US/08/973,124
                                                                                                                                                                                                                                                                                                                                                                                                                   304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07/931,473
                                                                                                                                                                                                                                                                    3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08/117,991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07/964,624
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 24.6; DB 1; Length 71; Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NEX 34-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19; Indels
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Gaps

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PCT-US96-08014-78/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 78, Application PC/TUS9608014 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/45/
FILING DATE: 02-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: I
                                                                                                                                  COMPUTER: IBM compatible OPERATING SYSTEM: MS-DOS SOFTWARE: WOOTDPETFECT 6.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (303) 793-34
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: LARRY GOLD; NEBOJSA JANJIC; STEVEN RINGQUIST; APPLICANT: PAGRATIS; PENELOPE J. TOOTHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3958 gaacatgtgattaggctgtattcaaagcagatcaccattgcgatcaacgtcctcc 4012
                 APPLICATION NUMBER:
                                                                                              CLASSIFICATION:
                                                                                                                FILING DATE
                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                   CITY: Englewood
                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
PRIOR APPLICATION UNBER: 08/465,591
PRIOR APPLICATION NUMBER: 05-JUNE-1995
                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           56 GAAGACGTAAGTAAGTATTAAAAACATACCCCCCCTGGCCGCATCGTCCTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 20-MARCH-19: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 07-JUNE PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 07-JUNE PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            / Match 0.5%; Score 24.6; DB 4; Local Similarity 65.5%; Pred. No. 1.2e+03; nes 36; Conservative 0; Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                     Colorado
                                                                                                                                                                                                                                                                                          8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Barry
                                                                                                                                                                                                                                                      USA
                                                                                                                                                                                                                                                                                                         Swanson & Bratschun, L.L.C.
02-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UMBER: 08/479,783
07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MARCH-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           793-3433
                                                                                                                                                                                                                                                                                                                                                    KEROTINOCYTE GROWTH FACTOR (hKGF)
                                                                                                                                                                                                                                                                                                                                                                  HIGH AFFINITY OLIGONUCLEOTIDE LIGANDS TO TRANSFORMING GROWTH FACTOR (TGF), PLATELET-DERIVED GROWTH FACTOR (PDGF) AND HUMAN
          08/458,424
                                                        08/458,423
                                                                                                                       PCT/US96/08014
                                                                                                                                                                                                                                                                                                                                                 304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19; Indels 0; Gaps
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TOPOLOGY: 11;
MOLECULE TYPE:
PCT-US96-08014-78
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO: 78:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5,
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/100,664A
PILING DATE: 19-JUN-1998
FILING DATE: 19-JUN-1998
CLASSIFICATION: 435
ATTORNBY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REGISTRATION NUMBER: 600-1-221
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: YOUNG, MICHAEL W.
APPLICANT: KLOSS, BRIAN
APPLICANT: BLAU, JUSTIN
APPLICANT: PRICE, JEFFREY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/465,594
FILING DATE: 05-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 05-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/465
FILING DATE: 05-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/475
APPLICATION NUMBER: 08/475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/618
FILING DATE: 20-MARCH-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/47:
FILING DATE: 07-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 07-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/479,783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: A NOVEL CLOCK GENE AND METHODS OF USE TITLE OF INVENTION: THEREOF NUMBER OF SEQUENCES: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Barry J. Swanson REGISTRATION NUMBER: 33,215
                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                            ZIP: 0/001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 0.5%; Score 24.6; DB 5; Length 71; Local Similarity 65.5%; Pred. No. 1.2e+03; Local Similarity 65.5%; Mismatches 19; Indels es 36; Conservative 0; Mismatches 19;
                                                                                                                                                                                                                                                                 ZIP: 07601
                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , Application US/09100664A 6057129
                                                                                                                                                                                                                                                                                                                      Hackensack
                                                                                                                                                                                                                                                                                                    New Jersey
                                                                                                                                                                                                                                                                                                                                               411 Hackensack Avenue, 4th Floor
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US-09-100-664A-5
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Best Local S
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TELEX: 133521
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 72 base pairs
; TOPOLOGY: linear; MOLECULE TYPE: DNAUS-08-400-256-9
                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC PC DOS MS-DOS
OPERATING SYSTEM: PC PC DOS MS-DOS
SOFTWARE: PatcentIn Release #1.0, Version #1.25
SOFTWARE: PatcentIn Release #1.0, Version #1.25
CORRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/400,256
FILING DATE: 03-MAR-1995
CILASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Ellas J. 778
DESCREMENTANCE MINDREP: 33.778
                                                                                                         REGISTRATION NUMBER: 33.728
REGISTRATION NUMBER: 3985
REFERENCE/DOCKET NUMBER: 3985
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
TELEPHONE: 212-878-9655
TELEPAX: 212-878-9655
INFORMATION FOR SEQ ID NO: 9:
SEQUIENCE CHARRCTERISTICS:
LENGTH: 100 base pairs
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MOLECULE TYPE: CI
HYPOTHETICAL: NO
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Andersen, Asser Storin APPLICANT: Markussen, Jan TITLE OF INVENTION: ACYLATED INSULIN NUMBER OF SEQUENCES: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 405 Lexington Avenue, 04 Un CITY: New York STATE: 10174-6401 STATE: 10174-6401 COMPOTTER READABLE FORM: COMPOTTER READABLE: Floppy disk MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 57504970 No. 5750497disk of No. 5750497th America, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 0.5%;
Local Similarity 59.2%;
Les 42; Conservative
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                                                                LENGTH: 100 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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5. 5750497
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Halstrom, John
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US-08-975-365-9
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TELEPAX: 212-878-9655
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
ENGTH: 100 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9, Application US/08975365
Patent No. 6011007
                 4529 tatttacataaagactgtgctgagaagcag 4558
                                                                                 4469 aagaatgaagoctagtagaatatggacttggaaaattetettaateactactgtatgtaa 4528
                                                                                                                                                                                                                                 MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIDIM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
37 CAATTGGAAAACTACTGTGGTTAGACGCAG 8
                                                    97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTGTAC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Elias T
                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DALL.
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/400,256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                  . Match 0.5%; Score 24.4; DB 3; Length 100; Local Similarity 54.4%; Pred. No. 1.7e+03; Les 49; Conservative 0; Mismatches 41; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Lambiris, Elias J. REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 0
FILING DATE: 03-MAR-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: United States of America ZIP: 10174-6401
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                                                                                                                                                                                                                                                                                           100 base pairs
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Andersen, Asser Sloth
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Pred. No. 1.7e+03;
0; Mismatches 41; Indels
                                                                                                                    0; Gaps
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ICCATION: 2.88
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US-08-182-175A-54
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US-08-182-175A-54
                                                                 1120 gagtatgagggtgatgacacagaaggagaattaaaaggagttggaaagtagtacggatgag 1179
                          1180 agtgaagaagaa 1191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 54, Application US/08182175A Patent No. 5559223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/743,(
FILING DATE: 9 AUGUST 1991
ATTORNEY/AGENT INFORMATION:

NAME: Linda Axamethy Floyd
REGISTRATION NUMBER: 33,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Saverio Ca
APPLICANT: Sharon J.
APPLICANT: Janet A. F
                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
ORIGINAL SOURCE:
STRAIN: E. co.
                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
                                           10 GAAGATGAAGGCGATGGAGGAGAAGATGAAGTGGATGGAGGAGAAGATGAAGTGGATGGA 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 97 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: BB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (302) 992-4929
70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OFFWARE: Microsoft Word, 4.0
CURRENT APPLICATION DATA:
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TELEFAX: 835420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Synthetic Storage Proteins with Defined Structure Containin NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                           NAME/KEY:
                                                                                                                                                                                                                                                                                                                                CLONE:
                                                                                                                                                                                                                                                                                                                                                          CELL TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1007 Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                         42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
                                                                                                                                                                                                                                                                                                                              92-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19898
                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                        E. coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Delaware
                                                                                                                                                                                                                                                                                             CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (302) 892-7949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: E.I. du Pont de Nemours and Company
                                                                                                                                                                                                                                                                                                                                                         DH5 alpha
                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Janet A. Rice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Saverio Carl Falco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                  DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                  double
                                                                                                                                                                                                    /function= "synthetic storage protein /product= "protein" /gene= "ssp" /standard_name= "5.11.11.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07/743,006
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/08/182,175A
                                                                                                                 0;
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                                                                                                                   Mismatches
                                                                                                              30; Indels
                                                                                                            0
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OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 63
Patent No.
GENERAL II
                            ğ
                                                            Дb
 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: E. I. DU PONT
APPLICANT: COMPANY
APPLICANT: COMPANY
TITLE OF INVENTION: CHIN
TITLE OF INVENTION: METE
TITLE OF INVENTION: AND
TITLE OF INVENTION: AND
TITLE OF INVENTION: OF:
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                      Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: FLOPF DISK

COMPUTER: IDM PC COMPATIBLE

COMPUTER: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: MICROSOFT WORD VERSION 2.0C

CURRENT APPLICATION DATA:

CURRENT APPLICATION NUMBER: US/08/474,633A

APPLICATION NUMBER: US/08/474,633A

FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
ATTORNEY/AGENT INF
                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                        NAME/KEY: CDS
LOCATION: 2..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: E. I. DU PONT DE NEMOURS
ADDRESSEE: AND COMPANY
STREET: 1007 MARKET STREET
CITY: WILLINGTON
STATE: DELAWARE
                                                                                                                                                                                                                                                                                                                                                                       IMMEDIATE
                                                                             1120 gagtatgagggtgatgacacagaaggagaattaaaaggagttggaaagtagtacggatgag 1179
                            1180 agtgaagaagaa 1191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                              CLONE: 92-2
                                                                                                                                                                                                                                                                                                                                                                                    STRAIN: E. coli
CELL TYPE: DH5 alpha
                                                          10 GAAGATGAAGGCGATGGAGGAGAAGATGAAGTGGATGGAGGAGAAGATGAAGTGGATGGA 69
70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3, Application US/08474633A 5773691
                                                                                                                       h 0.5%; Score 24; DB 1;
Similarity 58.3%; Pred. No. 2.2e+03;
42; Conservative 0; Mismatches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DU PONT DE NEMOURS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OF THE SEEDS OF PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CHIMERIC GENES AND
METHODS FOR INCREASING
INCREASING THE LYSINE
AND THREONINE CONTENT
                                                                                                                                                                                                                  /function="synthetic storage protein product= "protein" /product= "ssp" /standard_name="5.11.11.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BB-1037-C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INCREASING
                                                                                                                                   30; Indels
                                                                                                                                                             Length 97;
                                                                                                                                   0; Gaps
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RESULT 14
PCT-US92-06412-54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy Disk

MEDIUM TYPE: Floppy Disk

COMPUTER: MacIntosh System, 6.0

OPERATING SYSTEM: Macintosh System, 6.0

SOFTWARE: Microsoft Word, 4.0

CURRENT APPLICATION NATA:

APPLICATION NUMBER: PCT/US92/06412

FILING DATE: 19920807

FILING DATE: 19920807

FILING DATE: 19920807

FILING DATE: 9 AUGUST 1991

ATTORNEY AGENT INFORMATION:

PREFERENCE DOCKET NUMBER: BB-1031

REFERENCE DOCKET NUMBER: BB-1031

REFERENCE JOCCKET NUMBER: BB-1031

FELECOMMUNICATION INFORMATION:

TELEPHONE: (302) 992-7949

TELEFAX: (302) 992-7949

TELEFAX: (302) 992-7949

INFORMATION FOR SED ID NO: 54:

SEQUENCE CHARACTERISTICS:

LENGTH: 97 base pairs

STENENTFINESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 54, Application PC/TUS9206412
GENERAL INFORMATION:
APPLICANT: Saverio Carl Falco
                                                                                                                                                                                                                                                                  PCT-US92-06412-54
                                                                                                                                                                                                   Query Match
                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Saverio Carl Falco
APPLICANT: Sharon J. Keeler
APPLICANT: Sharon J. Keeler
TITLE OF INVENTION: Synthetic Storage Proteins with Defined Structure Containin
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
CLONE: 92-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
ORIGINAL SOURCE:
STRAIN: E. CO
                                                                                                                                                                                                                                                                            NAME/KEY: CDS
LOCATION: 2..88
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                      1180 agtgaagaagaa 1191
|| ||||| |||
                                                                                                     1120 gagtatgagggtgatgacacagaaggagaattaaaggagttggaaagtagtacggatgag 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1007 Mar)
CITY: Wilmington
STATE: Delaware
                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 19898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN: E. coli
CELL TYPE: DH5 alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                         10 GANGATGAAGGCGATGGAGGAGAAGATGAAGTGGATGGAGGAGAAGATGAAGTGGATGGA 69
70 AGAGAAGATGAA 81
                                                                                                                                                                            Local
                                                                                                                                                        tch 0.5%; Score 24; DB 5; Length 97; al Similarity 58.3%; Pred. No. 2.2e+03; al 2; Conservative 0; Mismatches 30; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2: E.I. du Pont de Nemours and Company 1007 Market Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                double
                                                                                                                                                                                                                                                                                       /function= "synthetic storage protein product= "protein" | gene= "ssp" | standard_name= "5.11.11.5"
                                                                                                                                                                               0;
                                                                                                                                                                                 Gaps
                                                                                                                                                                                    0
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CONTRACT OF A

ZIP: 10174-6401

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.30

APPLICATION NUMBER: US/08/672,158A

FILING DATE: 27-JUN-1996

APPLICATION NUMBER: 34,086

CLASSIFICATION NUMBER: 34,086

REFERENCE/DOCKET NUMBER: 34,086

REFERENCE/DOCKET NUMBER: 34,086

REFERENCE/DOCKET NUMBER: 3655.000-US

TELEPHONE: 212-867-0123

INFORMATION FOR SEQ ID NO: 8:

SEQUENCE CHARACTERISTICS:

LENGTH: 83 base pairs

TYPE: nucleic acid

STRANDEDNESS: single

MOLECULE TYPE: CDNA

US-08-672-158A-8
Search completed: May 29, 2002, 12:56:56 Job time: 9665 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 15
US-08-672-158A-8
                                                                                                                                                                              Matches
                                                                                                                                                                                                    Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8, Application US/08672158A Patent No. 5770371 GENERAL INFORMATION:
                                                                                  ADDRESSEE: No. 57703710 No. 5770371disk of No. 5770371th America, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NERAL INFORMATION:
APPLICANT: Sheryl Thompson
TITLE OF INVENTION: Modification of Cryptic Splice Sites In
NUMBER OF SEQUENCES: 21
                                                                                                                                                             y Match
0.5%; Score 23.8; DB 1; Length 83;
Local Similarity 72.1%; Pred. No. 2.3e+03;
hes 31; Conservative 0; Mismatches 12; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 405 Lex
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          405 Lexington Avenue, 64th Floor
                                                                                                                                                               0;
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Run on:

Title:

Sequence: perfect score:

Scoring table:

Searched:

Database :

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Result No.

Score

39.

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6 8 9 10 11 11 12 13 13 14 15

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OM nucleic - nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Minimum DB seq length: 0
Maximum DB seq length: 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-676-436-3
4990
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Maximum Match 100%
Listing first 45 summaries
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em_estpl:*
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95
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10 BI322285
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10 TB5845
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AA490364
AI767078
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BE573242
BF101686
AA267457
AA946634
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AV551027
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AA490364 aa44c10.r
AI767078 wi92g07.x
BJ063432 BJ063432
                                                                                                                              AI360096 qy89d09.x
BI322285 kx19a08.y
AA154655 mq61c08.r
T85845 yd61d10.r1
AI054386 qi76e03.x
C53310 C53310 Yuj1
BE573342 601333139
BE573342 601333139
                AA267457 mg90a03.r

AA946634 oq38f11.s

R93104 EST000038 S

R40354 yf81a05.s1

AV551027 AV551027
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMMENT
                                                                                                                                                                                                      FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JOURNAL
                                                                                                                                                                                      source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95 bp mRNA linear ES:
ua58f03.rl Soares_thymus_2NbMT Mus musculus cDNA clone
INAGE:1361691 5', mRNA sequence.
aA990067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Mus.

Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.

Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Muridae; Mus.

1 (bases 1 to 95)

2 (bases 1 to 95)

3 (bases 1 to 95)

3 (bases 1 to 95)

4 (bases 1 to 95)

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7 (bases 1 to 95)

8 (bases 1 to 95)

7 (bases 1 to 95)

8 (bases 1 to 95)

9 (bases 1 to 95)

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1 (bases 9 to 96)

2 (bases 9 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Marra M/Mouse EST Project
WashO-HHMI Mouse EST Project
WashIngton University School of MedicineP
WashIngton University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. LC
                                                                                                                                                                                                                                                                                                      Email: mouseest@watson.wustl.edu
Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                            Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The Washu-HHMI Mouse EST Project
                                                                                                                                                                                                                             seq primer: -28ml3 rev2 ET from Amersham High quality sequence stop: 82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Mus musculus"
/strain="057BL/6J"
/strain="157BL/6J"
/db_xref="taxon:1090"
/clone="IMAGE:1361691"
/clone_1ib="Soares_thymus_2NDMT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                     /sex="male"
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AII035450

AII340694

AII340694187

AII43956

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AV532477

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AV532479
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TA330B01P
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B1858071 603384887
BH127076 G-10924.f
AA865746 0h41h11.s
A1035450 ub47b5.r
A1035450 ub47b5.r
A1035450 ub47b5.r
A1035450 ub47b5.x
A1035450 ub47b5.x
A104597 0h86b08.s
A1494187 mc67e06.x
A1142956 0z58d01.x
A1086378 0z44c01.x
A1086378 0z44c01.x
A1086378 0z44c01.x
A1086378 0z44c01.x
A108638 0z45201.x
A108638 0z45201.x
A108638 0z45201.x
A108638 0z45201.x
A166505 A166509 A1665095
A166505 A1665095
A1665095 A1665095
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AI339161 qt14h01.x
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AL492149 T. brucei
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REFERENCE
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AA490364
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                          TIMAGE Consortium (info@image.lln1.gov) for further information. Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq_primer: -28ml3 rev2 ET from Amersham
                                                                                                                                                                                                                                                                                                                       Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                              Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lavy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                   WashU-Merck EST Project 1997
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA490364 70 bp mRNA linear EST 08-AUG-1997 aa44c10.r1 Soares_NhHMPU_S1 Homo sapiens cDNA clone IMAGE:823794 5' similar to SW:NIM1_NEUCR P48479 PROTEIN KINASE NIM-1;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                        clone is available royalty free through LLNE; contact the
                                                                                                                                                                                                                                                                                                         314 286 1800
314 286 1810
                                                                                                                                                                                                                                                                                    est@watson.wustl.edu
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/note="Organ: mixed (see below); Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site_1: Not I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="Pooled human melanocyte, fetal heart, and pregnant uterus"
                                                                                         /clone="IMAGE:823794"
/clone_lib="Soares_NhHMPu_S1"
                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                'lab_host="DH10B"
                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="Thymus"
/dev_stage="4 weeks"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%;
70.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 39.8; DB 9;
Pred. No. 24;
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                                                                                                                                                                                                                                                                                                                                                                                                 Insert Length: , J. J. Seq primer: -40UP from Gibco Seq primer: -6001000 from Fibration (1998) 70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                      www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 785 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nlh.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 71)
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w192907.xl NCI_CGAP_Kidl2 Homo sapiens cDNA clone IMAGE:2400828 3',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AI767078.1 GI:5233587
             /note="Organ: Kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_CGAP_Kid5 was prepared, and ss circles were made in vitro. Following HAP hybridization reaction. The driver was PCR-amplified cDNAs has used as tracer in a subtractive from a pool of 5,000 clones made from the same library (cloneIDs 1323912-132881, 1471368-1472903 and 1492104-149225). Subtraction by Bento Soares and M. Fatima Bonaldo. "Subtraction by Bento Soares and M.
Fatima Bonaldo. "
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                                                                                                                                                                                                                                   /clone_lib="NCI_CGAP_Kid12"
/tissue_type="2 pooled tumors (clear cell type)"
                                                                                                                                                                                                                                                                                           /clone="IMAGE: 2400828"
                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.6%; Score 29; DB 9;
71.7%; Pred. No. 1.4e+04;
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RESULT 4
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AI360096/c
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      135 tttgaaagaaaaatgaatgcaccaaatcagcctccacataaagacactggaaaaacagt 194
                                                                                                                                                                                                                                                                                                                                                                                195 gga 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%;
Similarity 64.2%;
13; Conservative
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BJ063432 NIBB Mochil normalized Xenopus tailbud library Xenopus
BJ063432 NIBB Mochil normalized Xenopus tailbud library Xenopus
Laevis CDNA clone XL076el3 5', mRNA sequence.
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1 (bases 1 to 83)
1 (chases 1 to 83)
1 (this of the state of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Tadasu Shin i
Center For Genetic Resource Information
Rational Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
Tel: 81-559-81-6856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Expressed genes in X. laevis embryo unpublished (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: tshini@genes.nig.ac.jp.
Location/Qualifiers
   AI360096 83 bp mRNA linear EST 16-FEB-1999 qy89d09.x1 NCI_CGAP_Brn25 Homo sapiens CDNA clone IMAGE:2019185 3', mRNA sequence.
AI360096 1 GI:4111717 EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.6%; Score 28; DB 10;
Similarity 63.5%; Pred. No. 2.6e+04;
Strain Conservative 0; Mismatches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Xenopus laevis"
/db_xref="taxon:8355"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="stage 25"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="NIBB Mochii normalized Xenopus tailbud
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="whole_embryo"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Score 28.6; DB 9; Length 71; pred. No. 1.7e+04; O; Mismatches 24; Indels
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BI322285
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NCJ/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological National Institut
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 27.8; DB 9; Length 83; Ch 12 Similarity 59.5%; Pred. No. 3e+04; Pred. No. 3e+04; O; Mismatches 32; Indels
                                                                                                                                                                                                                                                                                                   94 bp mRNA linear EST 30-JUL-2001
BI322285 mRNA linear EST 30-JUL-2001
kx19a08.y3 Parastrongyloides trichosuri cDNA 5', mRNA sequence.
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Panagrolaimoidea; Strongyloididae; Parastrongyloides. 1 (bases 1 to 94) 1 (bases 1 to 94) 1 (bases 1, to 94) McCarter, J., Clifton, S., Chiapelli, B., Pape, D., Martin, J., Wylle, T., McCarter, J., Clifton, S., Chiapelli, B., Pape, D., Martin, B., Bowers, Y., Dante, M., Marra, M., Hillier, L., Kucaba, T., Theising, B., Bowers, Y.
                                                                                                                                                   parastrongyloides trichosuri
parastrongyloides trichosuri
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                                                                                                AUTHORS
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l (bases 1 to 100)
Marra M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                             AA154655

AA154655

IQO bp mRNA linear EST 19-FEB-1

Similar to SW:KCC4_MOUSE PO8414 CALCIUM/CALMODULIN-DEPENDENT

PROTEIN KINASE TYPE IV CATALYTIC CHAIN; mRNA sequence.
                                                                                                                                                                                                                  AA154655.1 GI:1726483
                                                                                                                                                                                   nouse mouse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The library was constructed by Brandi Chiapelli and Dr. James McCarter (bchiapel@watson.wustl.edu & jmccarte@watson.wustl.edu) at Washington University, St. Louis. DNA Sequencing by: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer:
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The vector to vector length is 95
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Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The Washington Univ. Nematode EST Project, 1999 Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: McCarter JP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gibbons,M., Ritter,E., Bennett,J., Franklin,C., Tsagareishvili,R. Ronko,I., Kennedy,S., Maguire,L., Beck,C., Underwood,K., Steptoe M., Allen,M., Person,B., Swaller,T., Harvey,N., Schurk,R., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and
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314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="vector: pAMP1 (Gibco); Site_1: NotI; Site_2: SalI; The library was constructed by Brandi Chiapelli and Dr. Dames McCarter at Washington University, St. Louis. The PCR based library using Dynabead oliopedT priming (Dynal). SMART PCR cDNA Synthesis Kit from Clontech. Directionally provided by Dr. Warwick Grant of AgRearch, New Zealand (warwick grant@agresearch.co.nz)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    est@watson.wustl.edu
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Parastrongyloides trichosuri FL pAMP1 v1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="parastrongyloides trichosuri"
/db_xref="taxon:131310"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 94;
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4037 acattaaaggtgccaatatcttcctta 4063
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                                                                                                                                                                                                                                                                                                                                          27 ATCTAAAAGCTAGCAATATTCTCTATA 1
yd61d10.rl Soares fetal liver spleen INFLS Homo sapiens cDNA cl
                                                                                                                                       Homo sapiens
                                                                                                                                                                                  T85845.1 GI:714197
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                                                                                                                                                                                                            HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Possible reversed clone: similarity on wrong strand
Seq primer: -28M13 rev2 from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The WashU-HHMI Mouse EST Project Unpublished (1996) Contact: Marra M/Mouse EST Project WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMAGE Consortium (info@image.linl.gov)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicinep
4444 Forest Park Parkway, Box 8501, St. Ld
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - Oligo(dT) primer [5, 37]; double-stranded cDNA was ligated to Eco RI adaptors and Eco RI sites of the modified pT7T3 vector. RNA rounds of normalization, and was constructed by Bento Soares and M.Fatima Bonaldo."

24 a 11 c 20 g 45 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /dev_stage="4 weeks"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Soares_thymus_2NbMT"
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/clone="IMAGE:583214"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /strain="C57BL/6J"
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57.5%;
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Pred. No. 3.2e+04;
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Insert Size: 715
Insert Size: 715
High quality sequence starts: 1 High quality sequence stops: 1
High quality sequence starts: 1 High quality sequence royalty-free
Source: IMAGE Consortium, LLML This clone is available royalty-free
through LLML; contact the IMAGE Consortium (info@image.lln1.gov)
through LLML; contact the IMAGE Consortium (info@image.lln1.gov)
for further information. Trace considered overall poor quality
for further unique; 715
Std Error: 0.00
seq primer: M13RP1
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4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
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Contact: Wilson RK
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                            AIO54386 98 bp mRNA linear EST 27-JUL-1998 qi76e03.x1 NCI_CGAP_Ov26 Homo sapiens cDNA clone IMAGE:1862428 3', mRNA sequence.
AIO54386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           n 0.5%;
Similarity 54.4%;
Tumor Gene Index Unpublished (1997) Contact: Robert Strausberg, Ph.D. Email: cgapbs-remail.nih.gov Tissue Procurement: Monica Brown, M.D., Elise Kohn, M.D., Michael R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 98)
                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                               AI054386.1 GI:3322173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /dev_stage="20 week-post conception fetus"
/dev_stage="20 week-post conception fetus"
/dev_stage="DH10B (ampicillin resistant)"
/note="Organ: Liver and Spleen; Vector: pT7T3D (Pharmacia)
/note="Organ: Liver and Spleen; Vector: pT7T3D (Pharmacia)
/note="Organ: Liver and Spleen; Vector: page 1. Site_2: Eco RI;
/with a modified polylinker; Site_1: Page I oligo(dT) primer
ist strand cDNA was primed with a Page I oligo(dT) primer
ist stranded cDNA was ligated to Eco Ri adaptors
double-stranded cDNA was ligated to Eco Ri adaptors
(Pharmacia), digested with Page I and cloned into the Page I
and Eco RI sites of the modified pT7T3 vector. Library
went through one round of normalization. Library
vent through one round of normalization. Library
constructed by Bento Soares and M.Fatima Bonaldo."

8 a 20 c 20 g 33 t 10 others
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:9606"
/clone="IMAGE:112723"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="GDB:468340"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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FEATURES
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Best Local Similarity
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA Sequencing by: Washington University Genome Sequencing Center
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/pbrp/image/image.html
www-bio.llnl.gov/pbrp/image/image.html
seq primer: -40ml3 fwd. ET from Amersham.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100 bp mRNA linear EST 11-SEP-
C53310 Yuji kohara unpublished cDNA Caenorhabditis elegans cDNA
Clone yk259b3 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C53310.1 GI:2391067
                                                                                                                                                                                                                                                                                                                                                               Kohara,Y., Motohashi,T., Tabara,H., Watanabe,H., Sugimoto,A., Sano,M., Miyata,A. and Nishigaki,A.
Expression map of the C.elegans genome
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Caenorhabditis elegans
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                                                                                                                                                                                                             Genome Biology Lab.

Genome Biology Lab.
National Institute of Genetics
Yata 1111, Mishima, Shizuoka 411, Japan
Yata 181-559-81-6855
Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                             Unpublished (1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1862428"
/clone_lib="NCI_CGAP_Ov26"
                                                                                                                                                                         ykohara@lab.nig.ac.JP.
Location/Qualifiers
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/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                Yuji Kohara
                                                             /organism="Caenorhabditis elegans"
/organism="Caenorhabditis elegans"
/strain="CB1489 him 8 (e1489)"
/db_xref="taxon:6239"
/clone="yk259b3"
/sex="hermaphrodite, male"
/tissue_type="whole animal"
                                      /clone_lib="Yuji Kohara unpublished cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%; Score 27.4; DB 9;
62.3%; Pred. No. 4.1e+04;
ative 0; Mismatches 26;
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BASE COUNT
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                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                     41;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-remail.nih.gov
Tissue Procurement: Jeffrey Green M.D.
cDNA Library Preparation: Life Tenchnologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
plate. ITAMOGOO
                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                       Plate: LLAM8933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        house mouse.
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                                                                                                                                                                      /note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies. Investigator providing samples: Jeffrey Green, M.D., NIH"
                                                                                                                                                                                                                                                              /Clone_lib="NCI_CGAP_Mam6"
/Sex="female, virgin"
/tissue_type="infiltrating ductal carcinoma"
/dev_stage="5 months"
                                                                                                                                                                                                                                                         /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:3710769"
                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/strain="FVB/N"
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7 c 18 g
                                                                                   0.5%; Score 26.6; DB 10; Length 81; 63.1%; Pred. No. 6.1e+04; Indels
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Mus musculus
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IMAGE:720652 5', mRNA sequence.
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              house mouse.
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5 ATAAGGCTGCTTACTAGAGTTTAATGCAAATGTTCATAGCTCATTTCTTTTTTTATCATG 64
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM9177 row: d column: 06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Muses 1 to 82)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /strain="FVB/N"
/db_xref="taxon:10090"
/clone="mAGE:3980885"
/clone_lib="NGI_CGAP_Mam1"
/tissue_type="tumor, biopsy sample"
/dev_stage="10 months, virgin"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH"

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Pred. No. 6.1e+04;
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REFERENCE
AUTHORS
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AA946634/c
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WashU-HHMI Mouse EST Project
Washington University School of MedicineP
Washington Park Parkway, Box 8501, St. Louis, MO 63108
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1810
Fax: 314 286 1810
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1 (bases 1 to 87)

Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Gelsel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, Mylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, Mylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, Mylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, Mylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, Mylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, M., Mylie, T., Lennon, G., Soares, B., Wilson, R. and Dietrockers, M., Dietrich, N., Dubuque, T., Dietrich, N., Dubuque, T., Soares, B., Wilson, R., and Dietrockers, M., Dietrich, N., Dubuque, T., Soares, B., Wilson, R., and Dietrockers, M., Dietrich, N., Dubuque, T., Dietrich, N., Dubuque, T., Soares, B., Wilson, R., M., Dietrich, N., Dietrich, N
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This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 95)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA946634
oq38f11.s1 NCI_CGAP_Kid5 Homo
mRNA sequence.
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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/lab_host="DH10B"
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/dev_stage="4 weeks"
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sapiens cDNA clone IMAGE:1588653 3',
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Inpublished (1997)

AL Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Contact: Robert Strausberg, Ph.D.

Emmil: cgapbs-remail.nih.gov

Fissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.

Fissue Procurement: Christopher Moskaluk, M.D., Ph.D.

Chan Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CLONA Library Preparation: M. Bento Soares, Ph.D.

CLONA Library Preparation: M. Bento Soares, Ph.D.

CLONA Library Preparation: M. Bento Soares, Ph.D.

Clone distribution: NCI-CGAP clone distribution information can be clone distribution: M. A.G.E. Consortium/LLNL at:

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/inage/lmage/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lmgde/lm
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                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Platyhelminthes; Trematoda; Digenea; Strigeldida; Schistosomatoidea; Schistosomatidae; Schistosoma.

1 (bases 1 to 95)
1 (bases 1 to 95)
2 (bases 1 to 95)
3 (bases 1 to 95)
5 (bases 1 to 95)
6 (bases 1 to 95)
7 (bases 1 to 95)
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1 (bases 1 to 95)
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EST000038 S. mansoni cDNA Schistosoma mansoni cDNA clone
SMTBADAMS0038SK 5' end, mRNA sequence.
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                                                                                       TBRI Biochemistry
Theodor Bilharz Research Institute
Theodor Bilharz Research Institute
Tmbabb, p.O.Box 12411, Giza, Egypt
Tel: 202 31218276
Fax: 202 3121167
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/tissue_type="2 pooled tumors (clear cell type)"
/tissue_type="2 pooled tumors (page: type)"
/note="Organ: kidney; Vector: pT7T3D-Pac (pharmac
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ORIGIN
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                                                                                                                                                                                                       Query Match 0.5%; Score 26.4; DB 10; Length 95; Best Local Similarity 59.2%; Pred. No. 7.3e+04; Matches 45; Conservative 0; Mismatches 31; Indels 0; Gaps
                                                                                                                                   701 tcaaagtcgactatgg 716
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: sk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: M-Saber@FRCU.EUN.EG
                                                                                                                                                                                                                                                                                                                                   31 a
                                                                                                                                                                                                                                                                                                     /organism="Schistosoma mansoni"
/organism="Egyptian"
/db_xref="taxon:6183"
/clone=!bs"S. mansoni cDNA"
/lab_host="E. coli XL Bluel"
/note="Vector: pBluescript II SK+; Site_1: EcoR1; Site_2: CDNA was constructed and cloned simultaneously using wector priming with the pBluescript II SK+ vector. cDNA vector priming with the pBluescript II SK+ vector. cDNA vector to the Xhol site."

15 c 10 g 39 t
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Post-processing: Minimum Match 0%
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                                                                                                                                                                                                                                                             Result
No.
                                                                                                                           000
                                                                                                                                                                                                                                                                                                                                                                                        pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                   Score
N_Geneseq_032802:*
1: /SIDS1/gcgdata/gc
2: /SIDS1/gcgdata/gc
3: /SIDS1/gcgdata/gc
4: /SIDS1/gcgdata/gc
5: /SIDS1/gcgdata/gc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NICHESSEY_OGGATA/geneseq/geneseqn-embl/NA1980.DAT:

2: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1981.DAT:

3: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1982.DAT:

3: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1984.DAT:

5: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1984.DAT:

5: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1985.DAT:

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7: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1985.DAT:

8: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1988.DAT:

9: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1989.DAT:

10: SIDS1/gcgdata/geneseq/geneseqn-embl/NA1989.DAT:

11: SIDS1/gcgdata/geneseq/geneseqn-embl/NA199.DAT:

12: SIDS1/gcgdata/geneseq/geneseqn-embl/NA199.DAT:

13: SIDS1/gcgdata/geneseq/geneseqn-embl/NA199.DAT:

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14: SIDS1/gcgdata/geneseq/geneseqn-embl/NA199.DAT:

15: SIDS1/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:

16: SIDS1/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:

17: SIDS1/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:

18: SIDS1/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:

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16: SIDS1/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:

17: SIDS1/gcgdata/geneseq/geneseqn-embl/NA2
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Listing first 45 summaries
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00000000
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                  Mercuric ion reduc
primer 307-3398 fo
GFP Leu(CTG)5 forw
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Human brain natriu
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	AAK23782 AAK49875 AAI26980 AAI55798 AAI1529		AAA92397 AAAH51645	ABA69691 AAC12392 AAS06930 AAC11499 AAC11499 AAC15541 AAC152344 AAX23469 AAX25128 ABA66615 ABA66615
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ALIGNMENTS

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RESULT
AAT50934
AAT50934;
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                                                                                                                                                                        p53 binding protein; Mdmx; tumour suppressor; cancer; ss
                                                                                                                                                                                        Mouse p53-recognition clone 1 5' sequence
                                                                                                                                                                                                          07-APR-1997 (first entry)
                                                                                                                                          W09641875-A1.
                                                                                                                                                           Mus sp.
                                                                                                         13-JUN-1996;
                                                                                                                          27-DEC-1996.
      Nucleic acid encoding p53-binding protein - which has homology to mouse Mdm2, for use in cancer research
                                                                 (INTR-) INTROGENE BV.
(UYLE-) RIJKSUNIV LEIDEN.
                                                                                         13-JUN-1995;
                                 WPI; 1997-065462/06.
                                                  Jochemsen A,
                                                                                                                                                                                                                                                            ۳,
                                                   Shvarts A,
                                                                                            95EP-0201565
                                                                                                            96WO-NL00239.
                                                      Van Der Eb AJ;
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RESULT 2
AAC19445/c
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          The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively total human RNAs or polya+ RNAs derived from 30 different tissues. EST of the mRNA because they are often obtained from oligo-dr primed cDNA derived from the 5' ends of mRNAs and even in those cases where longer 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be in diagnostic. Gene have been obtained, the full 5' UTR is rarely included. used to obtain full length cDNAs and genomic DNAs. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be in diagnostic. Gene therefore be the diagnostic. Gene therefore be the diagnostic. Gene therefore be and constituted from the full suited for manactives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                               Claim 1; SEQ ID 23520; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                    New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                      WPI; 2000-500381/45.
                                                                                                                                                                                                                                                                                                                                                   Dumas Milne Edwards J,
                                                                                                                                                                                                                                                                                                                                                                                     (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                           21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted protein 5' EST, SEQ ID NO: 23520.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-OCT-2000 (first entry)
diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC19445;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC19445 standard; cDNA; 92 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequences of the 5' region (AAT50934) and 3' region (AAT50935) were detd. for cDNA clone 1, isolated from a mouse 16-day embryo cDNA library using tumour suppressor p53 as probe. The 5' and 3' regions (AAT50936-37) of another clone were also detd. These submitted to several databases. A third clone (see also AAT50933) coded for a p53 binding protein designated Mdmx (AAW10206).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 78 BP; 33 A; 9 C; 22 G; 10 T; 4 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 2a; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 tgvnghagacaaaa 74
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ilarity 58.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              99US-0122487
                                                                                                                                                                                                                                                                                                                                         Duclert A, Giordano J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 27.2; DB 10;
Pred. No. 1e+03;
"""matches 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 78;
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4469 aagaatgaagoottagtagaatatggacttggaaaattotottaatcactactgtatgtaa 4528

0; Mismatches Score 26; Pred. No.

2.7e+03;

40;

Indels

0;

Gaps

DB 16; Length 100;

97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTGTAC 38

Matches Query Match Best Local

50;

Local

0.5%; Similarity 55.6%; Conservative

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                      AAQ86399 and AAQ86400 are a pair of primers for the PCR amplification of the Asp3 B-chain/Ala21 A-chain human insulin derivative DNA. The two subunits of the derivative are covalently connected via disulphide bonds between Cys residues A7/B7 and A20/B19. The derivative, which may be present as a zinc ion complex, can be used as a fast action treatment for diabetes.
Sequence 100 BP; 29 A; 22 C; 23 G; 26 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                    Example 1; Page 30; 100pp; English.
                                                                                                                                                         Acylated insulin deriv. which may be present as a Zinc ion complex - is used to treat diabetes and is rapid acting.
                                                                                                                                                                                                                  WPI; 1995-131314/17.
                                                                                                                                                                                                                                                 Markussen J;
                                                                                                                                                                                                                                                               Andersen As,
                                                                                                                                                                                                                                                                                         (NOVO ) NOVO-NORDISK AS
                                                                                                                                                                                                                                                                                                                            02-FEB-1994;
                                                                                                                                                                                                                                                                                                                                       17-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                    16-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                  23-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                               W09507931-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human insulin derivative; A-Chain; diabetes; Zinc ion complex Ala21 PCR primer #128; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human insulin derivative Ala21 A-chain PCR primer #128.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ86400 standard; DNA; 100 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ86400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 92 BP; 12 A; 9 C; 15 G; 47 T; 9 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression and secretion vectors.
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                                                                                                                                                                                                                                                          Halstrom JB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                    94US-0190829
                                                                                                                                                                                                                                                                                                                                       93DK-0001044
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57.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 26.6; DB 21;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                    Jonassen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.7e+03
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RESULT 4
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ID AAH4868/C
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AC AAH486
AC A
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH48689 standard; DNA; 60 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human G-protein subunit Gbeta3 intron 9 DNA fragment #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-OCT-2001 (first entry)
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21-FEB-2000; 2000DE-1007587.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-JUN-2000; 2000DE-1030945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel use for sequence variants of the human G-protein beta3 subunit to predict physiological and pathological pathways in the human body. The invention may be used to detect, detect pathways into for, give a prognosis of, or treat a variety of diseases predisposition for, give a prognosis of, or treat a variety of diseases and other heart or circulation diseases, disbectes, psychiatric diseases and other heart or circulation diseases, disbectes, psychiatric diseases, immunological diseases, asthma, psoriasis, Crohn's disease, colities ulcerosa, transplant rejection, HIV or Hepatitis B or C. This sequence represents a fragment of the human G-protein Gbeta3 subunit intron 9 which is used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Use of sequence variants of the beta3 subunit of human G protein to predict, detect, and determine drug targets for diseases including heart, immunological and psychiatric diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SIFF/) SIFFERT W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-490007/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 3; 14pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 60 BP; 6 A; 11 C; 6 G; 37 T; 0 other;
AAA71594;
                                                                                                                                                                                                                                                           507 ctcagtctcaaagaaaaagacagggagcaaagagggacaagaaaatacgtctgg 560
                                                               AAA71594 standard; DNA; 70 BP.
                                                                                                                                                                                                                     Local Similarity
                                                                                                                               υ
                                                                                                                                                                                                                                                                                                                                                        36;
                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                          0.5%;
                                                                                                                                                                                                                                                                                                                                                        Score 25.2; DB 22; Length pred. No. 3.4e+03; O; Mismatches 18; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 60;
                                                                                                                                                                                                                                                                                                                                                                                      0;
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SØ SØ
                                                                                                                                                                                                                                                                     B
                                                                                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brain natriuretic peptide; BNP; primer; human; detection; epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human brain natriuretic peptide N-terminal primer Pro2hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method to detect N-terminal pro-brain natrituretic peptide (BNP) in a sample using two antibodies, that natrituretic peptide (BNP) in a sample using two antibodies, that recognize different epitopes of the N-terminal pro-BNP. The method can be used for diagnosing heart insufficiency. The recombinant N-terminal pro-BNP is useful as a standard in a method to detect N-terminal pro-BNP and also for production of antibodies against N-terminal pro-BNP this sequence represents a primer used in the cloning of recombinant rhis sequence represents a primer used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-1999;
12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200045176-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identification of N-terminal pro-brain natriuretic peptide (pro-BNP) using two antibodies that recognize different epitopes of the N-terminal pro-BNP, useful for diagnosis of heart insufficiency -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-499359/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HOFF ) ROCHE DIAGNOSTICS GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JAN-2000; 2000WO-EP00602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Karl J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 32; 35pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 70 BP; 10 A; 19 C; 21 G; 20 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                  1778 agggcggcctgctgatgaagcagtactaccagttcatgctgcaggaggttctggaggact 1837
                                                                                                                                                                                                                                                                                               1838 tg 1839
                                                                                                                                                                                     AAT22086 standard; cDNA to mRNA; 89 BP.
                        Gene signature; messenger RNA; mRNA; relative abundance; frequency human; cloning; mapping; non-biased library; diagnosis; detection; cell typing; abnormal cell function; ss.
                                                                                                                          02-AUG-1996 (first entry)
                                                                                       Human gene signature HUMGS03631.
Homo sapiens.
                                                                                                                                                                                                                                                                     68 tg 69
                                                                                                                                                                                                                                                                                                                               8 agggaggtotgttcaacctgcagttcggacagittaccctgcaggtggttacgctgttcc 67
                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                       σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lill H, Stahl P, Krueger K, Borgya A,
                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99DE-1003489.
99DE-1011044.
                                                                                                                                                                                                                                                                                                                                                                                                                      0.5%;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                        Score 25.2; DB 21;
Pred. No. 3.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gallusser A;
                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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W09514772-A1

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AAT92244
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                                                                                                                                                                                                                                              Query Match
     16-SEP-1997.
                  US5668294-A
                                      Synthetic
                                                toxic heavy metal; transgetransposon; Tn21; PCR; ss
                                                                      Mercuric ion reductase merA PCR mutagenesis primer 307-339S.
                                                          Metal ion resistance; mercuric ion reductase; merA; mercury;
                               Transposon Tn21
                                                                                          19-JAN-1998
                                                                                                                  AAT92244 standard; DNA; 99 BP
                                                                                                         AAT92244;
                                                                                                                                                                   4545 gt 4546
                                                                                                                                                                                 Sequence 89 BP; 39 A; 8 C; 15 G; 27 T; 0 other;
                                                                                                                                                                         Claim 1; Page 1039; 2245pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                              Identifying gene signatures in 3'-directed human cDNA library - e.g. for diagnosis of abnormal cell function, by preparing cDNA that reflects relative abundance of corresp. mRNA in specific human
                                                                                                                                                      73 gt 74
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-206931/27.
                                                                                                                                                                                                                                                                                                                                                                                                                            Matsubara K, Okubo K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-1995.
                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                           (OKUB/) OKUBO
                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-NOV-1994;
                                                                                                                                   7
                                                                                                                                                                                                                                                                                                                                                                                                                                               (MATS/) MATSUBARA K.
                                                                                                                                                                                                            39;
                                                                                                                                                                                                                 Similarity
                                                                                        (first entry)
                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            93JP-0355504.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94WO-JP01916
                                                   transgenic plant; mutagenic; bacterial;
                                                                                                                                                                                                               0.5%;
                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                            Score 25.2; DB 16;
Pred. No. 4.4e+03;
                                                                                                                                                                                                       23;
                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                  Length 89;
                                                                                                                                                                                                     0;
                                                                                                                                                                                                   Gaps
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PI
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DR
XX
PT
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XX
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SO
SO
SO
SX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                       WPI; 1999-579950/49.
                                                                         (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                            21-APR-1995;
                                                                                                                                        19-JUN-1997;
                                                                                                                                                                                                                                                        aquatic environment; bioremediation; contamination; PCR; amplification.
                                                                                                                                                                                                                                                                      Organometal; resistance; transcription; translation; regulation; merA; transposon; Tn2l; merB; mercuric ion reductase; organomercury lyase; ss; transgenic plant; detoxification; metal compound; soil; sediment; primer
                                                                                                                                                                                                  US5965796-A.
                                                                                                                                                                      12-OCT-1999
                                                                                                                                                                                                                              Transposon In21.
                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                       Primer 307-339S for Tn21 merA gene.
                                                                                                                                                                                                                                                                                                                                                                     24-DEC-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         AAZ28594;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3086 ttgattttctacaagcaattgaacctgcctttat 3119
                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ28594 standard; DNA; 99 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3026 atgteetgaetaaatgtgagagtggtagaggtacaagacecaggtgggegaeteaaggat 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a PCR mutagenesis primer used to mutagenise the merA coding sequence derived from transposon Tn21, to adapt it for plant expressibility. The nucleic acid molecule produced is useful for producing transgenic plants that are resistant to toxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 99 BP; 29 A; 21 C; 25 G; 24 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           heavy metals (especially mercury) and so can be used remediate and revegetate contaminated areas. The bacterial mera gene is derived the transposon Tn21, which was originally isolated from the incompatibility Group IncFII resistance plasmid NRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Column 43-44; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        producing transgenic plants resistant to toxic heavy metals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62 gtgagcatactcaagcaagccaagttgcctatat 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA encoding mercuric ion reductase in plant-expressible form - for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-470112/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Meagher RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 aagacccagctataggtgaagctgttactgctgcatttcgcatggaaggcattgaagtgc 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
                                                 Meagher RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                    97US-0878957.
                                                                                                         95US-0427097.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Summers AO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95US-0427097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-0427097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.5%;
54.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 25.2; DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ŏ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.7e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Metal resistance sequences for producing transgenic plants

Page 5

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The invention relates to a nucleic acid molecule comprising a coding code sequence for an organometal resistance protein, which is operably control of a nucleic acid molecule comprising a coding colinked downstream of and under the regulatory control of a great plant-expressible transcription and translation regulatory sequence. Concern the companion of the Ecoli merB gene merA encodes a mercuric ion concern the molecular concern the Ecoli merB gene merA encodes a mercuric conference of the concern the Ecoli merB gene merA encodes a mercuric conference are useful for producing transgenic plants which are capable conference are useful for producing transgenic plants are able to remove or conference are useful for producing transgenic plants are able to remove or conference are useful compounds such as methyl mercury and ionic mercury from conference and aquatic environments providing an efficient conference and advantace environments providing an efficient conference are expressed in the sequence represents a pCR primer used to amplify the Tn21 merA conference are expressed in the conference of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Column 17-18; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 99 BP; 29 A; 21 C; 25 G; 24 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3026 atgtcctgactaaatgtgagagtggtagaggtacaaggacccaggtgggcgactcaaggat 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3086 ttgattttctacaagcaattgaacctgcctttat 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA73946 standard; DNA; 54 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                        Green fluorescent protein; GFP; reporter gene; codon utilisa translational efficiency; protein abundance; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GFP Leu(CTG)5 forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-2000
                                                                                                                                                                                                                                                                                                                                                          WO200042215-A1.
                                                                                                                                                                                                                                                                                                                                                                                        synthetic.
                                                                                                                                                                                                                                                                                                                        20-JUL-2000
                                                                                                                                                                                                                                                      08-JAN-1999;
                                                                                                                                                                                                                                                                                      07-JAN-2000; 2000WO-AU00008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62 gtgagcatactcaagcaagccaagttgcctatat 95
The present sequence is a primer used to generate a synthetic gfp gene by
                                                               Determining translational efficiency of codons in cells, comprintroducing synthetic constructs with reporter genes fused in tandem repeats of the codon, and measuring expression -
                                                                                                                                   WPI; 2000-499118/44.
                                                                                                                                                                       Zhou J,
                             Example 1; Page 183; 190pp; English.
                                                                                                                                                                                                                    (UYQU ) UNIV QUEENSLAND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 aagacccagctataggtgaagctgttactgctgcatttcgcatggaaggcattgaagtgc 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  similarity 54.3%;
                                                                                                                                                                       Frazer IH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                          99AU-0008078
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 25.2; DB 20;
Pred. No. 4.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         codon utilisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 99;
                                                                                                     frame
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cc codon followed by a stretch of five identical codons was fused in frame cc codon followed by a stretch of five identical codons was fused in frame cc immediately upstream of a gfp coding sequence to form the synthetic gene. CC The amplified fragment was cloned into the mammalian expression vector cc cc promas, which contains SV40 ori and the CMV promoter, and was used in a cc promas, which construct was introduced into COS-1 cells and expression of cc the reporter protein (green fluorescent protein) was measured. A series crepater protein (green fluorescent protein) was measured. A series co f 4 gfp reporter constructs was made in which the gfp gene is preceded cc in frame by a tandem repeat of 5 identical codons. The series covers the cc entire set of isoaccepting codon triplets. Codons with a higher cc translational efficiency than their corresponding synonymous codons can cc preferred codons of a polynucleotide so that there is higher protein cc preferred codons of a polynucleotide so that there is higher protein cc expression within undifferentiated epithelial cells such as COS-1 cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABA69691,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 54 BP; 9 A; 14 C; 20 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2732 ccagtcagccggtcatcgccaaagctttgcagcagctgaagaatgatgc 2780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABA69691 standard; DNA; 91 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA69691;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; foetal liver; gene expression; single exon nucleic acid probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human foetal liver single exon nucleic acid probe #17996.
                                                                                                                                                                                                                                                                                                                                                      04-FEB-2000; 2000US-0180312.

26-MAY-2000; 2000US-0207456.

30-JUN-2000; 2000US-0608408.

03-AUG-2000; 2000US-0632366.

03-AUG-2000; 2000US-0234687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-JAN-2001; 2001WO-US00669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200157277-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     53 CCAGTGAACAGTTCCTCGCCCTTGCTCAGCAGCAGCAGCAGCATGGTAC 5
The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid probe of the invention.
                                                                                                                                                                                                                                                                                                                     27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                          Claim 4; SEQ ID NO 17996; 639pp + sequence listing; English.
                                                                                                                                                         Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver \dot{\,}
                                                                                                                                                                                                                 WPI; 2001-483447/52.
                                                                                                                                                                                                                                                    Penn
                                                                                                                                                                                                                                                                                   (MOLE-) MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               34;
                                                                                                                                                                                                                                                      SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                      Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.5%;
                                                                                                                                                                                                                                                           Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 25; DB 21;
Pred. No. 3.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                               Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 54;
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ş
of the mRNA because they are often obtained from oligo-dy primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences cDNA sequences have been obtained, the full 5' UTR is rarely included. Set to obtain full length cDNAs with intact 5' ends and can therefore be in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
                                                                                                                                                                                                The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID 16467; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dumas Milne Edwards J, Duclert A, Giordano J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-500381/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-FEB-2000; 2000EP-0200610.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted protein 5' EST, SEQ ID NO: 16467.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC12392;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC12392 standard; cDNA; 77 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3815 acgaattgaaaatattcgaaggcatcaaacaccccaatctggttcggtattttggtgtgg 3874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3875 agctc 3879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 91 BP; 28 A; 19 C; 13 G; 31 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format dir.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   67 AAGAAATTATTATGATGAAAGACTGTAAACACCCAAATATTGTTGCTTATTTTGGAAGCT 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7 ATCTC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5' EST; expressed sequence tag; secreted protein; cDNA isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 25; DB 22,
Pred. No. 5.1e+03;
"" matches 25;
                                                                                                                                                                                                                                              ESTs were prepared from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      format directly
                                                                                                                                                                                                                            tissues. EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
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Sequence 77 BP; 27 A; 14 C; 15 G; 21 T; 0 other;

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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QΥ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 В
          genome, or antigenome combined with one or more heterologous gene(s) or more heterologous pathogen(s) to form a chimeric genome or more heterologous pathogen(s) to form a chimeric genome or antigenome or more heterologous pathogen(s) to form a chimeric genome or antigenome. The chimeric PIV is useful in an attenuated vaccine to elicit an immune response against one or more virus(es) selected from human PIV1 (HPIV1), response against HPIV3, measles or RSV. An immunospecific composition an immune response against HPIV3 and the second chimeric PIV elicits immune response against HPIV3 and the second chimeric PIV elicits an second chimeric PIVs elicit an immune response against HPIV1 or HPIV2, and where both the first and pathogen. Chimeric HPIV3, HPIV1 and HPIV2 are useful as vaccines to
prevent measles and upper or lower respiratory tract
                                                                                                                                                                                                                                                                                                              The sequence is described in an invention relating to novel infectious chimeric parainfluenza viruses (PIVS). The virus comprises a major nucleocapsid protein (N), a nucleocapsid phosphoprotein (P), a large polymerase protein (L), and a partial or complete PIV vector background
                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence for reverse PCR primer for respiratory syncytial virus (RSV) A G gene insert is used with the forward PCR primer (AAS06929) in the construction of bovine/humanpIV3-RSV chimeric viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 150; 305pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated infectious chimeric parainfluenza virus (PIV), useful in an attenuated vaccine to elicits an immune response against one or more virus(es) selected from human PIV1 (HPIV1), HPIV2 and HPIV3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-356173/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murphy BR, Collins PL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-DEC-1999;
10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-DEC-2000; 2000WO-US33293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200142445-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleocapsid phosphoprotein; large polymerase; attenuated vaccine; human PIV1; HPIV1; HPIV2; HPIV3; RSV; pathogen; measles; PCR primer; respiratory syncytial virus; respiratory tract infection; bovine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Infectious chimeric parainfluenza virus; antigenic determinant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse PCR primer for RSV A G gene insert.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAS06930 standard; DNA; 92 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS06930;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14 TATCTCAG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 74 GACTGTGTAGGTGCATTTGTTTAAAAAATATAACTCCCCATTGCCTCTGGGATAAAGTCT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 60.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0459062.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schmidt AC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ,
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Pred. No. 5.2e+03;
.^. Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Durbin AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Skiadopoulos MH;
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AAC11499
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Best Local
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                                                                                                                                                            The present sequence is one of a large number of 5' ESTS derived from CC mRNAs encoding secreted proteins. No ORF has yet been conclusively controlled within the present sequence. The 5' ESTS were prepared from CC total human RNAs or polyA+ RNAs derived from 30 different tissues. EST CC total human RNAs or polyA+ RNAs derived from 30 different tissues. EST CC of the mRNA because they are often obtained from oligo-off primed cDNA cC libraries. Such ESTs are not well suited for isolating cDNA sequences CC libraries. Such ESTs are not well suited for isolating cDNA sequences CC cDNA sequences of mRNAs and even in those cases where longer CC cDNA sequences have been obtained, the full 5' UPR is rarely included. CC cDNA sequences have been obtained, the full 5' UPR is rarely included. S' ESTS are derived from mRNAs with intact 5' ends and can therefore be considered to obtain full length cDNAs and genomic DNAs. 5' ESTS are also used to obtain upstream regulatory sequences and to design they are used to obtain upstream regulatory sequences and to design or expression and secretion vectors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      particularly in young children
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 92 BP; 20 A; 12 C; 26 G; 34 T; 0 other;
                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted protein 5' EST, SEQ ID NO: 15574.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC11499;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC11499 standard; cDNA; 94 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dumas Milne Edwards J, Duclert A, Giordano J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80 TTCATCTCCACCCAACACACCACGCCAGTAGTAAAAAAAGTAAGAAAAACTTAGGATTAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45 tacototocococoagoacacotogacagatgaaacgcatgtcaaccaaacatcagaggaa 104
                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID 15574; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-500381/45
1379 gaccatttgtagacaaagcactgaagcagatggggttaagaaagttaattttaagacttc 1438
                                                                                                                                  Sequence 94 BP; 46 A; 9 C; 14 G; 25 T; 0 other;
                                                       Local Sil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                       0.5%;
Similarity 54.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome mapping; ss.
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                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                           Score 24.8; DB 21; Length 94; pred. No. 6e+03;
                                                                   42; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                       0;
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10 CCTTCC 5

1655 tctgcc 1660

Matches Query Match

39;

Local

Similarity Conservative

0.5%;

1595 tgtcqtqqqaqqaqctqaaqqccatggatttaccttcattcqaacctqccttcctagttc 1654

Mismatches

6e+03;

25;

Indels

0;

Gaps

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AAC15541,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1439 acaagctaatggatggttccttgcaaagggca 1470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC15541 standard; cDNA; 94 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted protein 5' EST, SEQ ID NO: 19616.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC15541;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-OCT-2000 (first entry)
                                                                                                             The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST total human RNAs or polyA+ RNAs derived from 30 different tissues. EST the mRNA because they are often obtained from oligo-dT primed cDNA of the mRNA because they are often obtained from oligo-dT primed cDNA of the mRNA sequences albraries. Such ESTs are not well suited for isolating cDNA sequences they are often the full 5' UTR is rarely included. cDNA sequences have been obtained, the full 5' UTR is rarely included. cDNA sequences have been obtained, the full 5' UTR is rarely included. cDNA sequences have been obtained, the full 5' UTR is rarely included used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used used to obtain upstream regulatory sequences and to design expression and serverior vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62 aagatttaatgtaggaaatattattaaaggca 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dumas Milne Edwards J, Duclert A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID 19616; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-500381/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 gaaaatataaaaatactaaccaaaagaagtctgaggtaactatggtaatattaaaactcc 61
                                                              Sequence 94 BP; 35 A; 10 C; 29 G; 16 T; 4 other;
                                                                                                    expression and secretion vectors.
Score 24.8;
Pred. No. 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Giordano J;
                          DB 21;
                          Length 94;
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AAT65208/c RESULT 15

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Matches
                                                                            Query Match
                                                                                                                           Sequence 71 BP; 15 A; 16 C; 22 G; 18 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-MAR-1996;
02-JUN-1995;
02-JUN-1995;
05-JUN-1995;
05-JUN-1995;
07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 15; Page 123; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identification of nucleic acid ligands to TGF-beta, PDGF and hKGF using SELEX, used in the diagnosis and treatment of proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-034387/03.
                  MATCH 0.5%;
Local Similarity 65.5%;
Local Similarity 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NEXS-) NEXSTAR PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-DEC-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Transforming; growth factor; beta-1; TGF-beta-1; binding ligand; identification; SELEX; anti-mitogenic; inhibition; cell; Systematic Evolution of Ligands by Exponential enrichment; epithelial; proliferation; diagnosis; treatment; fibroids;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9638579-A1.
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95US-0458423.
95US-0458424.
95US-0465591.
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95US-0479725.
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       Score 24.6; DB 18;
Pred. No. 5.7e+03;
0; Mismatches 19;
     Indels
                                                    Length 71;
0; Gaps
0,
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Search completed: May 29, Job time: 9833 sec

2002, 13:05:25

QΥ

3958 gaacatgtgattaggctgtattcaaagcagatcaccattgcgatcaacgtcctcc 4012

56 GAAGACGTAAGTAAGTATTAAAAAACATACCCCCCCTGGCCGCATCGTCCTCC 2

Mr.			
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Run on:

Scoring table: Sequence: Title: Perfect score:

Searched:

Minimum DB seq length: 0
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Database

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Result No.

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14: 15: 16: 17: 18: 19: 20: 21: 22: 22: 23: 24: 25: 25: 26: 27:

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10: 11: 12:

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                    pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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Match Length DB
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em_htg_inv:*
em_htg_other:*
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                                                        Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.

1 (bases 1 to 87)

1 (bases 1 to 87)

Haribabu,B. and Dottin,R.P.

Identification of a protein kinase multigene family of Dictyostelium discoideum: Molecular cloning and expression Dictyostelium discoideum: Molecular cloning and expression CDNA encoding a developmentally regulated protein kinase Proc. Natl. Acad. Sci. U.S.A. 88, 1115-1119 (1991)
                                                                                                                 protein kinase 4.
Dictyostelium discoideum (strain Ax-3) DNA.
Dictyostelium discoideum
                                                                                                                                        M59747.1 GI:167723
                                                                                                                                                                                                                               /organism="Dictyostelium discoideum"
/strain="AX-3"
/db_xref="taxon:44689"
<1...>87
                                           Location/Qualifiers
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AF050514
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AF050514 Human end
M28495 Trypanosoma
AJ130784 Mus muscu
AX326634 Sequence
AX326671 Sequence
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195107 Sequence 78
S76509 DIS8 (A) {h
L20950 Drosophila
AR007408 Sequence
U92173 Mus musculu
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AB048411 Homo sapi
AR014621 Sequence
BD010413 Chimeric
I26754 Sequence 54
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AR007403 Sequence
AF025984 Paralicht
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AR062856 Sequence
L31186 Human STS U
AX326603 Sequence
AX059602 Sequence
AF025983 Paralicht
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AR140801 Sequence
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M14473 Rabbit musc
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AX033159 Sequence
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AL033823 H. sapiens
M19461 Chicken c-m
U72578 Oryctolagus
AF174523 Bufo dana
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M32437 Rat/polyoma
AX240922 Sequence
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                                                                                                                                                                       INV 27-APR-1993
                                                                                                                                                                                                                                              Human pre-B
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Best Local Similarity 59.5%;
                                                                                                                                                                                                                                                                           4529 tatttacataaagactgtgctgagaagcag 4558
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            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectoidei; Paralichthyidae; Paralichthys.

1 (Dases 1 to 96)
Quattro, J.M. and Jones, W.J.
                                                                                    Paralichthys dentatus
                                                                                                   summer flounder.
                                                                                                                               AF025984.1 GI:2570847
                                                                                                                                                                                                                                                                                                                                                                                   50;
                                                                                                                                                           96 bp DNA linear VRT 30-OCT-PAILICHTHYS dentatus lactate dehydrogenase (LDHA) gene, allele
                                                                                                                                                                                             AF025984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent: US 5750497-A 4 12-MAY-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                           0.5%;
Similarity 55.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Havelund, S., Halstr
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22 c 23 g
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/protein_id="ARA33189.1"
/db_xref="GG:167724"
/translation="NLLIDQYGHIKLTDFGFAKRITENTKSMC"
a 12 c 14 g 25 t
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                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                  Score 26; DB 6; Length 100; Pred. No. 1.7e+05;
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Best Local Similarity
Matches 39; Conserv
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68 TG 69
                                            8 AGGGAGGTCTGTTCAACCTGCAGTTCGGACAGTTTACCCTGCAGGTGGTTACGCTGTTCC 67
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 GTCGGACAAATCCTAGACGATTCATTTACAAAACAGTGTGCAGATGAGTAATGAGTCCT 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 0.5%;
Local Similarity 64.4%;
                                                                                                                                                                                                                                                                                         GALLUSSER ANDREAS (DE); KARL JOHANN (DE)
STAHL PETER (DE); KRUEGER KERSTIN (DE);
ROCHE DIAGNOSTICS GMBH (DE)
                                                                                                                                                                                                                                                                                                                                Gallusser,A., Kari,J., Lill,H., Stahl,P., Krueger,K. and Borgya,A. wethod of identifying n-terminal probnp Patent: WO 0045176-A 03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Escherichia coli.
Escherichia coli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 (bases 1 to 96)
Quattro, J.M. and Jones, W.J.
Direct Submission
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                                                                                                                      Conservative
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                                                                                                                                                                                                                               /db_xref="taxon:562"
                                                                                                                                                                                                                                               organism="Escherichia coli"
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/allele="PdLDHA2"
                                                                                                                        0.5%; Score 25.2; DB 1; Length 70; 62.9%; Pred. No. 2.7e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Paralichthys dentatus"
/db_xref="taxon:66718"
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BORGYA ANNELIESE (DE) ;
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RESULT AX033159

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                      Local
                                                               62 GTGAGCATACTCAAGCAAGCCAAGTTGCCTATAT 95
                                                                                                                           2 AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTCGCATGGAAGGCATTGAAGTGC 61
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Sequence 3 from Patent W00045176.
AX033159
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1 (bases 1 to 70)
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Sequence 5 from patent US 5668294.
165769
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Meagher, R.B. and Summers, A.O.
Metal resistance sequences and transgenic plants
Patent: US 5668294-A 5 16-SEP-1997;
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                                                                                                                                                                                                          Similarity
                                                                                                                                                                                       0.5%; Score 25.2; DB 6; Length 99; 1larity 54.3%; Pred. No. 2.8e+05; Conservative 0; Mismatches 43; Indels
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/db_xref="taxon:562"
/db_x1 19 c 21 g 20 t
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1 21 c 25 g
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Query Match 0.5%;
Best Local Similarity 61.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       78 GTACT 82
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryotolagus.

1 (bases 1 to 83)
1 (bases 1 to 84)
1 (bases 1 to
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23 a 13 c 27 g
About 719 bp after segment 17
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18 of 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence 31 from Patent W00142445.
                                                                                                                                                                                                                                                                                                                                                       Murphy, B.R., Collins, P.L., Schmidt, A.C., Durbin, A.P., Skiadopoulos, M.H. and Tao, T. Use of recombinant parainfluenza viruses (plvs) as vectors to protect against infection and disease caused by piv and other human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                synthetic construct synthetic construct
                                                                                                                                                                                                                                         Patent: WO 0142445-A 31 14-JUN-2001; The Secretary of the Department of H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              artificial sequence.
1 (bases 1 to 92)
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/db_xref="taxon:9986"
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/organism="synthetic construct"
/ob_xref="taxon:32630"
/ob_ref="taxon:52630"
/note="Reverse primer for RSV A G gene insert"
12 c 26 g 34 t
                                                                                                                                                                                          cary of the Department of Health and Human Services (US)
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Pred. No. 3.1e+05;
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Score 24.8; DB 6;

Length 92;

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RESULT 1
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AR140801/c
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Best Local Similarity 65.5%;
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Best Local Similarity 65.5%;
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                                                                                                                                           3958 gaacatgtgattaggctgtattcaaagcagatcaccattgcgatcaacgtcctcc 4012
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                                                                                                                              56 GAAGACGTAAGTAAGTATTAAAAAACATACCCCCCTGGCCGCATCGTCCTCC 2
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                                     71 bp
Sequence 78 from patent US 5731424.
195107
                                                                                                                                                                                               36;
                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 71)
Toothman,P.J., Ringquist,S. and Gold,L.
High affinity TGF.beta. nucleic acid ligands
Patent: US 5731144-A 78 24-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 78 from patent US 5731144.
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Gold,L., Janjic,N. and Pagratis,N.
High affinity oligonucleotide ligands to growth factors
Patent: US 6207816-A 78 27-MAR-2001;
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Sequence 78 from patent US 6207816.
AR140801
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Pred. No. 3.9e+05;
0; Mismatches 19;
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Pred. No. 3.9e+05;
0; Mismatches 19; Indels
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                                                                              4914 tgtaatgcaaaaggctgattactgaaatttaagaaaaaggttcttttttcaataaatggt 4973
                                     4974 ttattt 4979
                                                                                                                                                                                                                                                                                                                gene
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12 TTATTT 7
                                                                                                                                                            y Match 0.5%;
Local Similarity 60.6%;
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Local Similarity 65.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                   entry [NCBI gibbsq 76509] from the original journal article. This sequence comes from Figure 2.
                                                                                                                                                                                                                                                                                                                                                                                                                          GenBank staff at the National Library of Medicine created this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monckton,D.G. and Jeffreys,A.J.
Minisatellite 'isoallele' discrimination in pseudohomozygotes by
single molecule PCR and variant repeat mapping
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1 (bases I to 71)
Toothman,P.J., Ringquist,S. and Gold,L.
High affinity TGF.beta. nucleic acid ligands and inhibitors
Patent: US 5731424-A 78 24-MAR-1998;
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/allele="A"
a 0 c
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/db_xref="taxon:9606"
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Pred. No. 3.9e+05;
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DROTRANSIN

98 bp

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37 CAATTGGAAAACTACTGTGGTTAGACGCAG

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4529 tatttacataaagactgtgctgagaagcag. 4558
                                                                                                                                                                                                                                               source
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                                                                                                     Match 0.5%;
Local Similarity 54.4%;
tes 49; Conservative
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1 (bases 1 to 98)
Engels, W.R. and Johnson-Schlitz, D.M.
Deletion white allele. Transposase-Induced derivative of white-hd80k17
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AR007408 100 bp
Sequence 9 from patent US 5750497.
AR007408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1993)
2 (bases 1 to 98)
3 (bases 1 to 98)
4 (bases 1 to 98)
5 (bases 1 to 98)
6 (bases 1 to 98)
7 (bases 1 to 98)
7 (conversion of insertions and prelement-induced interallelic gene conversion of insertions and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 insertion sequence.

Drosophila melanogaster (individual_isolate B20.3) DNA.

Drosophila melanogaster
                                                                                                                                                                                                                                                                               Acylated insulin Patent: US 5750497-A 9 12-MAY-1998;
                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 100) Havelund, S., Halstr.o slashed.m, J., Jonassen, I., Andersen, A. Sloth.
                                                                                                                                                                                                                                                                                                                                                                    Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                         Unknown.
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ilarity 56.1%;
Conservative
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1 23 c 22 g
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/isolate="B20.3"
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                                                                                                                       Score 24.4; DB 6; Length 100; Pred. No. 4.5e+05;
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REFERENCE
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U92173/c
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                                                                                        1513 gaatttcagatcccatgfggggttcagattatgfgcagt 1552
1513 gaatttccagatcccatgfggggttcagattatgfgcagt 1552
41 GTATTTCCAAGATACCCTGTCGGGATCACTGCAGGTGCAGT 2
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                                                                                                                                                                 30;
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T cell receptor diversity in CTLs specific for the CTT-1 and CTT-2
minor histocompatibility antigens
J. Immunol. 159 (6), 2606-2615 (1997)
97444147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rochester, MN 55905, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Johnston, S.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 51)
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/product="T cell receptor beta chain"
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/db_xref="GI:2894947"
/translation="YCTCSDPDRYSGNTLYF"
a 13 c 11 g 15 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus"
/strain="C57BL/6"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                   /note="minor CTT-1 antigen specific CDR3 region"
                                                                                                                                                                                                                                                                                                                                                                                                                    /cell_type="cytotoxic T lymphocyte"
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1. .51
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75.0%;
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Pred. No. 5.6e+05;
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